



1600

RAW SEQUENCE LISTING

DATE: 05/06/2003

PATENT APPLICATION: US/09/870,353A

TIME: 14:43:20

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Output Set: N:\CRF4\05062003\I870353A.raw

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3 <110> APPLICANT: Wang, Yan
4     Xi, Lei
5     Prosen, Dennis E.
6     MJ Bioworks, Inc.
8 <120> TITLE OF INVENTION: Improved Nucleic Acid Modifying Enzymes
10 <130> FILE REFERENCE: 020130-000111US
12 <140> CURRENT APPLICATION NUMBER: US 09/870,353A
13 <141> CURRENT FILING DATE: 2001-05-30
15 <150> PRIOR APPLICATION NUMBER: US 60/207,567
16 <151> PRIOR FILING DATE: 2000-05-26
18 <150> PRIOR APPLICATION NUMBER: US 09/640,958
19 <151> PRIOR FILING DATE: 2000-08-16
21 <160> NUMBER OF SEQ ID NOS: 34
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33     binding protein Ssod7d
35 <220> FEATURE:
36 <221> NAME/KEY: CDS
37 <222> LOCATION: (1)..(189)
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55     binding protein Ssod7d
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131 Lys Met Ile Ser Phe Thr Tyr Asp Glu Gly Gly Gly Lys Thr Gly Arg
132           35           40           45
133 Gly Ala Val Ser Glu Lys Asp Ala Pro Lys Glu Leu Leu Gln Met Leu
134           50           55           60
135 Glu Lys Gln Lys Lys Gly Gly Val Thr Ser Pro Lys Ala Leu Glu
136   65           70           75           80
137 Glu Ala Pro Trp Pro Pro Pro Glu Gly Ala Phe Val Gly Phe Val Leu
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144           130          135          140
145 Ala Leu Arg Glu Gly Leu Gly Leu Pro Pro Gly Asp Asp Pro Met Leu
146 145          150          155          160
147 Leu Ala Tyr Leu Leu Asp Pro Ser Asn Thr Thr Pro Glu Gly Val Ala
148           165          170          175
149 Arg Arg Tyr Gly Gly Glu Trp Thr Glu Glu Ala Gly Glu Arg Ala Ala
150           180          185          190
151 Leu Ser Glu Arg Leu Phe Ala Asn Leu Trp Gly Arg Leu Glu Gly Glu
152           195          200          205
153 Glu Arg Leu Leu Trp Leu Tyr Arg Glu Val Glu Arg Pro Leu Ser Ala
154           210          215          220
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156 225          230          235          240
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159 Ala Glu Val Phe Arg Leu Ala Gly His Pro Phe Asn Leu Asn Ser Arg
160           260          265          270
161 Asp Gln Leu Glu Arg Val Leu Phe Asp Glu Leu Gly Leu Pro Ala Ile
162           275          280          285
163 Gly Lys Thr Glu Lys Thr Gly Lys Arg Ser Thr Ser Ala Ala Val Leu
164           290          295          300
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172          355          360          365
173 Ile Pro Val Arg Thr Pro Leu Gly Gln Arg Ile Arg Arg Ala Phe Ile
174          370          375          380
175 Ala Glu Glu Gly Trp Leu Leu Val Ala Leu Asp Tyr Ser Gln Ile Glu
176 385          390          395          400
177 Leu Arg Val Leu Ala His Leu Ser Gly Asp Glu Asn Leu Ile Arg Val
178          405          410          415
179 Phe Gln Glu Gly Arg Asp Ile His Thr Glu Thr Ala Ser Trp Met Phe
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181 Gly Val Pro Arg Glu Ala Val Asp Pro Leu Met Arg Arg Ala Ala Lys
182          435          440          445
183 Thr Ile Asn Phe Gly Val Leu Tyr Gly Met Ser Ala His Arg Leu Ser
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185 Gln Glu Leu Ala Ile Pro Tyr Glu Glu Ala Gln Ala Phe Ile Glu Arg
186 465          470          475          480
187 Tyr Phe Gln Ser Phe Pro Lys Val Arg Ala Trp Ile Glu Lys Thr Leu
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189 Glu Glu Gly Arg Arg Arg Gly Tyr Val Glu Thr Leu Phe Gly Arg Arg
190          500          505          510
191 Arg Tyr Val Pro Asp Leu Glu Ala Arg Val Lys Ser Val Arg Glu Ala
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200          580          585          590
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216 Ssod7d/full-length Taq
218 <220> FEATURE:
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220 <222> LOCATION: (1)..(2763)

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